Fig. 2 illustrates retroviral vectors pBIB-KS 1,2,3 modified to contain a Kosak translation initiation site and stop codons (SEQ ID Nos: 305-310).

Please replace the paragraph beginning at page 26, line 9, with the following rewritten paragraph:

Fig. 6 illustrates the 5' and 3' primer sequences (SEQ ID Nos: 304, 36, 37, 38) designed from C. pneumoniae which were used to isolate the SWIB and S13 genes from C. pneumoniae.

## **REMARKS**

The enclosed electronic and paper copies of the Sequence Listing include no new matter that goes beyond the original application as filed, but are supplied to fulfill the requirements as outlined in the Communication from the Examiner. Furthermore, the above amendments, which merely direct the insertion of the Sequence Listing and insertion of sequence identifiers, include no matter that goes beyond the original application as filed. Applicant respectfully submits that the above-identified application is now in compliance with 37 C.F.R. §§ 1.821-1.825.

Attached hereto is a marked-up version of the changes made to the specification and claims by the current amendment. The first of the attached pages is captioned "Version with Markings to Show Changes Made."

> Respectfully submitted, Seed Intellectual Property Law Group PLLC

Jane E.R. Potter

Ŕegistration No. 33,332

JEP:mls **Enclosures:** 

Computer Diskette
Declaration Regarding Computer Diskette
Paper Copy of Sequence Listing
Copy of Notice to Comply
701 Fifth Avenue, Suite 6300

Seattle, Washington 98104-7092 Phone: (206) 622-4900

Fax: (206) 682-6031

Application No.

09/454,684

Docket No.

210121.469C4

## VERSION WITH MARKINGS TO SHOW CHANGES MADE

In the Specification:

Paragraph beginning at page 26, line 1, has been amended as follows:

Fig. 2 illustrates retroviral vectors pBIB-KS 1,2,3 modified to contain a Kosak translation initiation site and stop codons (SEQ ID Nos: 305-310).

Paragraph beginning at page 26, line 9, has been amended as follows:

Fig. 6 illustrates the 5' and 3' primer sequences (SEQ ID Nos: 304,36,37,38) designed from C. pneumoniae which were used to isolate the SWIB and S13 genes from C. pneumoniae.